

**IN THE SPECIFICATION**

Please amend the paragraph starting on page 14, line 26 to page 15, line 3 as follows:

By contrast, *aroB* in *P. multocida* is located immediately upstream of *dam* with only a 4 bp gap between the genes, which is similar to the arrangement in *H. influenzae*. Our promoter analysis detected a potential promoter upstream of *P. multocida dam* within the *aroB* coding sequence. The 5' end of this promoter was located 115 bp upstream of the *P. multocida dam* start codon and had the sequence TGGAAA-17 bp-TAGCGT-5 bp-G (SEQ ID NO: 7).

Please insert the attached paper copy of the "Sequence Listing" after the last page of the specification.